

Notice of References Cited	Application/Control No. 09/975,072	Applicant(s)/Patent Under Reexamination ROCH ET AL.
	Examiner Christopher Nichols, Ph.D.	Art Unit 1647

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*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
	A	US-2003/0103980	06-2003	Korc et al.	424/155.1
	B	US-2002/0098511	07-2002	Heichman et al.	435/7.1
	C	US-			
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	N	WO 02/23193 A1	03-2002	WIPO	BETH ISRAEL DEACONNESS...	G01N 33/53
	O	WO 00/23109	04-2000	WIPO	UC REGENTS	A61K 39/395
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	U	Rosen et al. (August 1993) "Thyroxine Interactions with Transthyretin: A Comparison of 10 Different Naturally Occurring Human Transthyretin Variants." Journal of Clinical Endocrinology and Metabolism 77(2): 370-374
	V	Smeland et al. (1997) "Binding of perlecan to transthyretin in vitro." Biochem. J. 326: 829-836
	W	Lans et al. (4 April 1994) "Different competition of thyroxine binding to transthyretin and thyroxine-binding globulin by hydroxy-PCBs, PCDDs, and PCDFs." European Journal of Pharmacology 270(2-3): 129-136
	X	Golabek et al. (19 May 1995) "Amyloid b binding proteins in vitro and in normal human cerebrospinal fluid." Neuroscience Letters 191(1-2): 79-82

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
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	U	Wang & Liew (1994) "The human BAT3 ortholog in rodents is predominantly and developmentally expressed in testis." Molecular and Cellular Biochemistry 136: 49-57
	V	Bork and Bairoch (October 1996) "Go hunting in sequence databases but watch out for the traps." Trends in Genetics 12(10): 425-427
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	X	Smith and Zhang (November 1997) "The challenges of genome sequence annotation or 'The devil is in the details'." Nature Biotechnology 15:1222-1223

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	V	Skolnick and Fetrow (2000) "From gene to protein structure and function: novel applications of computational approaches in the genomic era." Trends in Biotech. 18(1): 34-39, especially p. 36 at Box 2
	W	Bork (2000) "Powers and Pitfalls in Sequence Analysis: The 70% Hurdle." Genome Research 10:398-400
	X	Ngo et al. (2 March 1995) "The Protein Folding Problem and Tertiary Structure Prediction, Chapter 14: Computational Complexity Protein Structure Prediction, and the Levinthal Paradox" pp. 492-495

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